

FIG. 2

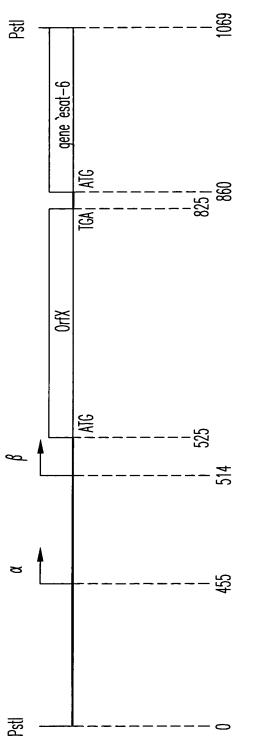


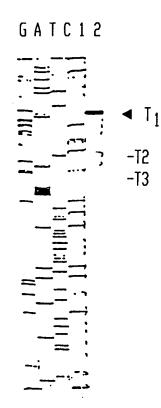
FIG. 3

G. FIG. SS SUBCLI <mark>ØBLION</mark> ET AL DOCKET #0660	413-3000 5-0X PCT SHEET 4 of 15	ussn 09/4	<b>0</b>
10	30	50	70
PstI . <u>CTGCAG</u> CAGGTGACGTC(	STTGTTCAGCCAGGTGGGCGGCA	CCGGCGGCGGCAACCCAG	CCGACGAGGAAGCCGCGCAGAT
90	110	130	150
GGGCCTGCTCGGCACCAC	STCCGCTGTCGAACCATCCGCTG	GCTGGTGGATCAGGCCCC	:AGCGCGGGCGCGGGCCTGCTGC
170	190	210	230
GCGCGGAGTCGCTACCTO	GGCGCAGGTGGGTCGTTGACCCG	CACGCCGCTGATGTCTCA	GCTGATCGAAAAGCCGGTTGCC
250	270	290	310
CCCTCGGTGATGCCGGC0	GGCTGTTGCCGGATCGTCGGTGA	CGGGTGGCGCCGCTCCGG	ITGGGTCCGGGAGCGATGGGCCA
330	350	370	390
GGGTTCGCAATCCGGCGC	GCTCCACCAGCCCGGGTCTGGTC	GCGCCGGCACCGCTCGCG	CAGGAGCGTGAAGAAGACGACG
410	430	450	470
AGGACGACTGGGACGAAG	GAGGACGACTGGTGAGCTCCCGT	+1 Mt AATGACAACAGACtTCCC	
490	510	1hp_	550
GCCAACATTTTGGCGAGG	. +1 Ms . GAAGGTAAAG <u>AGAGA</u> QAGTAGTCI	CAGCATGGCAGAGATGAA	GACCGATGCCGCTACCCTCGGG
570	RBS <b>→</b> 590	M A E M K 610	TDAATLG 630
CAGGAGGCAGGTAATTTC Q E A G N F 650	GAGCGGATCTCCGGCGACCTGA ERISGDLK 670	AAACCCAGATCGACCAGG T Q I D Q V 690	TGGAGTCGACGGCAGGTTCGTT ESTAGSL 710
GCAGGGCCAGTGGCGCGC Q G Q W R G 730	GCGCGGCGGGGGACGGCCGCCCAG A A G T A A Q A 750		CAAGAAGCAGCCAATAAGCAGA Q E A A N K Q K 790
AGCAGGAACTCGACGAGA Q E L D E I 810	TCTCGACGAATATTCGTCAGGC( S T N I R Q A 830	CGGCGTCCAATACTCGAG G V Q Y S R 850	GGCCGACGAGGAGCAGCAG A D E E Q Q Q esat-6
TAAAADDOTOOTOTOODOO	GGCTTCTGACCCGCTAATACGAA	AAAGAAACGGAGCAAAAA	CATGACAGAG

USSN 09/462,486

FIG. 4B

USSN 09/4 0



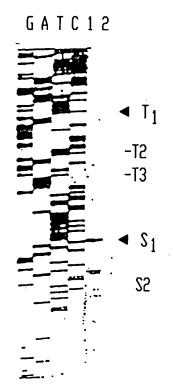


FIG. 5A

FIG.5C

AAGACGACGAGGACGACTGGGACGAAGAGGACGACTGGTGAGCTC

- 35

CCGTAATGACAACAGACTTCCCGGCCACCCGGGCCGGAAGACTTG

- 10

T 1

T 2

T 3

FIG. 5B



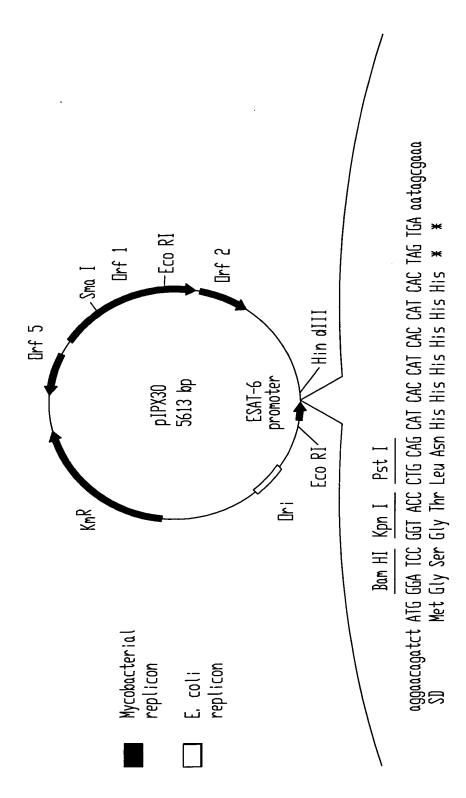
USSN 09/462,480



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cacgggatcgggggggttcgaccttccgtcggtctcgccct

ESAT-6 transcription terminator

FIG. 7

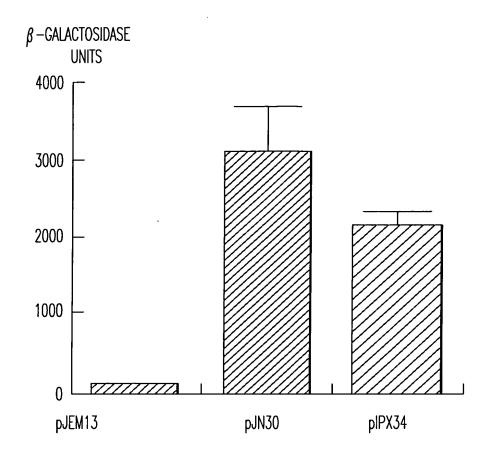
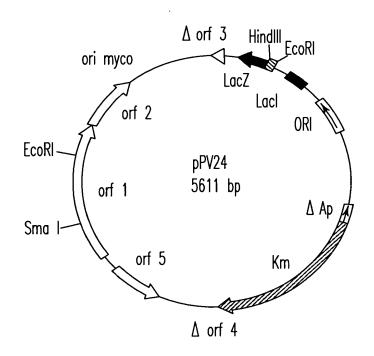


FIG.8

(His)

44

FIG. 9



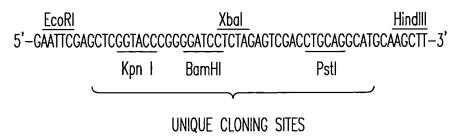
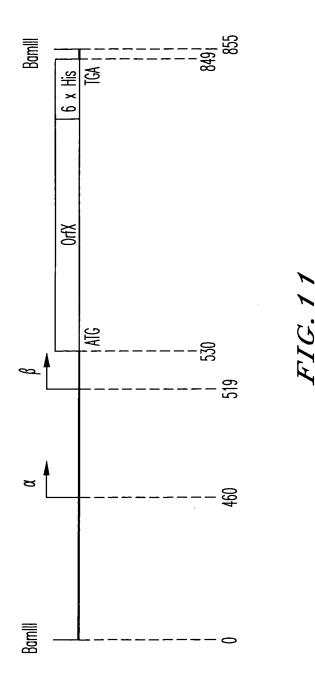
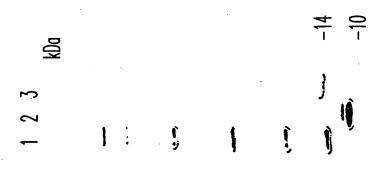
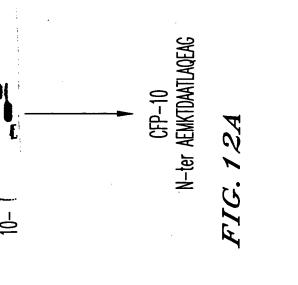


FIG. 10







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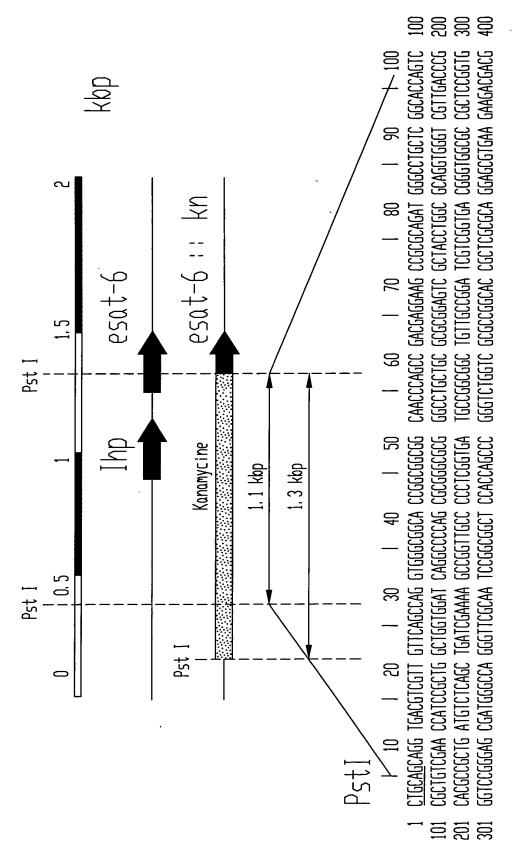


FIG. 134

1000 1069 99999 8 GCCGACCTGA TGCGCTTCCA GCAGCAGCAG TCGAGGCCGC GGCGAGGAA GCAGTGGAAT TTCGCGGGTA CCGACGAGGA GGCCGCCCAG GCCGCGGTGG GCGGATCTCC AAGCTCGCAG CGGCCTGGGG 8 **GCCAACATI** TACTCGAGGG GTAATTTCGA 8 CGGAAGACTT CGGCGTCCAA CAGGAGGCAG CGGCGGGGAC TGACAGAGCA GTCCCTGACC 2 ACCCTCGGG 909090999 JACTTCCCGG AGGGGAAGCA **GCTGAACAAC** TCGTCAGGC **AGCAAAAACA** 9 CTCCTTGACG GCAGGGCCAG TCGACGAATA **AAAGAAACGG** CGGCTACCGA 2 GAGATGAAGA CCGATGCCGC AATGACAACA CGACGAGATC CAGGTTCGTT GCTAATACGA CATTCATTCC TGGGACGCCA \$ GAGTCGACGG CAGCATGGCA CTTCTGACCC **ATGTCACGTC** CCAGCAAAAA AGCAGGAACT AGGACGACTG GGACGAAGAG GACGACTGGT 8 GAAAGTAGTC CGACCAGGTG AATAAGCAGA CGCAAATGGG ATCCAGGGAA **ACCAGGGTGT** ಜ **GGTAAAGAGA** AAACCCAGAT AGAAGCAGCC **JECAAGCGCA** 10 GCCTGTCCT CGGAGGCGT 

**16.13B** PSt[